RAW SEQUENCE LISTING DATE: 10/30/2003
PATENT APPLICATION: **US/09/924,103** TIME: 14:44:36

Input Set : **A:\18731055.app**
Output Set: **N:\CRF4\10302003\I924103.raw**

<110> APPLICANT: GOLDENBERG, DAVID M.
HANSEN, HANS J.
<120> TITLE OF INVENTION: IMMUNOTHERAPY FOR CHRONIC MYELOCYTIC LEUKEMIA
<130> FILE REFERENCE: 018733-1055
<140> CURRENT APPLICATION NUMBER: 09/924,103
<141> CURRENT FILING DATE: 2001-08-08
<160> NUMBER OF SEQ ID NOS: 4
<170> SOFTWARE: PatentIn Ver. 2.1

<210> SEQ ID NO: 1
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54 cttgtgaatg aagaagcaac cggacagttc catgtatacc gtgagtattt ccacatgacc 1920
55 tctgggtgtt gggggtcagt tctacttccc acatacggga ttgtcaggcc tgggttgtgc 1980
56 ctgtggccct ctctgcatta catcctgtat cagggtttgg acatttagtg caggacacac 2040
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61 <211> LENGTH: 141
62 <212> TYPE: PRT
63 <213> ORGANISM: Homo sapiens
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70              20                  25                  30
72 Thr Ala Lys Leu Thr Ile Glu Ser Thr Pro Phe Asn Val Ala Glu Gly
73          35                  40                  45
75 Lys Glu Val Leu Leu Leu Ala His Asn Leu Pro Gln Asn Arg Ile Gly
76      50                  55                  60
78 Tyr Ser Trp Tyr Lys Gly Glu Arg Val Asp Gly Asn Ser Leu Ile Val
79  65                  70                  75                  80
81 Gly Tyr Val Ile Gly Thr Gln Gln Ala Thr Pro Gly Pro Ala Tyr Ser
82                  85                  90                  95
84 Gly Arg Glu Thr Ile Tyr Pro Asn Ala Ser Leu Leu Ile Gln Asn Val
85             100                 105                 110
87 Thr Gln Asn Asp Thr Gly Phe Tyr Thr Leu Gln Val Ile Lys Ser Asp
88         115                 120                 125
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102 cgcccaccac tgctcagctc actattgaag ctgtgccatc caatgctgca gaggggaagg 180
103 aggttcttct acttgtccac aatctgcccc aggaccctcg tggctacaac tggtacaaag 240
104 gggaaacagt ggatgccaac cgtcgaatta taggatatgt aatatcaaat caacagatta 300
105 ccccagggcc tgcatacagc aatcgagaga caatataccc caatgcatcc ctgctgatgc 360
106 ggaacgtcac caaaaatgac acaggatcct acaccctaca agtcataaag ctaaatctta 420
107 tgagtgaaga agtaactggc cagttcagcg tacatccgga gactcccaag ccctccatct 480
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109 agactcagaa cacaacctac ctgtggtggg taaatggtca gagtctcccg gtcagtccca 600
110 ggctgcagct gtccaatggc aacaggaccc tcactctact cagtgtcaca aggaatgacg 660
111 taggacccta tgaatgtgaa atacagaacc cagcgagtgc aaacttcagt gacccagtca 720
112 ccctgaatgt cctctatggc ccagatgccc ccaccatttc cccttcagac acctattacc 780
113 atgcaggggt aaatctcaac ctctcctgcc atgcggcctc taatccaccc tcacagtatt 840
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117 tctcagctag agccactgtc agcatcatga ttggagtact ggccagggtg gctctgatat 1080
118 agtagctctg gtgtagtttc tgcatttcaa gaagactggc agacagttgt ttttattctt 1140
119 cctcaaagca tttgcaatca gctaccattc aaaattgctt cttcttcaag atttatggaa 1200
120 aatactctga cgagtactct tgaacacaag ttcctgataa ctttaagatc acgccactgg 1260
121 actgtctatg aacttgcaaa caggctgata cctttgtgaa gttgcccacc aaaacacaga 1320
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134 Gly Leu Leu Leu Thr Ala Ser Leu Phe Thr Phe Trp Asn Pro Pro Thr
135             20                  25                  30
137 Thr Ala Gln Leu Thr Ile Glu Ala Val Pro Ser Asn Ala Ala Glu Gly
138         35                  40                  45
140 Lys Glu Val Leu Leu Leu Val His Asn Leu Pro Gln Asp Pro Arg Gly
141     50                  55                  60
143 Tyr Asn Trp Tyr Lys Gly Glu Thr Val Asp Ala Asn Arg Arg Ile Ile
144 65                  70                  75                  80
146 Gly Tyr Val Ile Ser Asn Gln Gln Ile Thr Pro Gly Pro Ala Tyr Ser
147                 85                  90                  95
149 Asn Arg Glu Thr Ile Tyr Pro Asn Ala Ser Leu Leu Met Arg Asn Val
150             100                 105                 110
152 Thr Lys Asn Asp Thr Gly Ser Tyr Thr Leu Gln Val Ile Lys Leu Asn
153         115                 120                 125
155 Leu Met Ser Glu Glu Val Thr Gly Gln Phe Ser Val His Pro Glu Thr
156     130                 135                 140
158 Pro Lys Pro Ser Ile Ser Ser Asn Asn Ser Asn Pro Val Glu Asp Lys
159 145                 150                 155                 160
161 Asp Ala Val Ala Phe Thr Cys Glu Pro Glu Thr Gln Asn Thr Thr Tyr
162                 165                 170                 175
164 Leu Trp Trp Val Asn Gly Gln Ser Leu Pro Val Ser Pro Arg Leu Gln
165             180                 185                 190
167 Leu Ser Asn Gly Asn Arg Thr Leu Thr Leu Leu Ser Val Thr Arg Asn
168         195                 200                 205
170 Asp Val Gly Pro Tyr Glu Cys Glu Ile Gln Asn Pro Ala Ser Ala Asn
171     210                 215                 220
173 Phe Ser Asp Pro Val Thr Leu Asn Val Leu Tyr Gly Pro Asp Ala Pro
174 225                 230                 235                 240
176 Thr Ile Ser Pro Ser Asp Thr Tyr Tyr His Ala Gly Val Asn Leu Asn
177                 245                 250                 255
179 Leu Ser Cys His Ala Ala Ser Asn Pro Pro Ser Gln Tyr Ser Trp Ser
180             260                 265                 270
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183         275                 280                 285
185 Ile Thr Thr Lys Asn Ser Gly Ser Tyr Ala Cys His Thr Thr Asn Ser
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189 305                 310                 315                 320
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192                 325                 330                 335
194 Ser Ile Met Ile Gly Val Leu Ala Arg Val Ala Leu Ile
195             340                 345
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